

SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: (other than US) THE WALTER AND ELIZA HALL INSTITUTE OF MEDICAL RESEARCH, COUNCIL FOR THE QUEENSLAND INSTITUTE FOR MEDICAL RESEARCH LIMITED, UNIVERSITY OF MELBOURNE

(US only) Perry F BARTLETT, Andrew W BOYD, Mirella DOTTORI, Lynne HARTLEY, Mary GALEA, George PAXINOS, Mark POLIZZOTTO, Trevor KILPATRICK, Mark MURPHY, Frank KONTGEN, Jason COONAN, Ursula GREFERATH

(ii) TITLE OF INVENTION: A METHOD OF TREATMENT

(iii) NUMBER OF SEQUENCES: 6

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: DAVIES COLLISON CAVE

(B) STREET: 1 LITTLE COLLINS STREET

(C) CITY: MELBOURNE

(D) STATE: VICTORIA

(E) COUNTRY: AUSTRALIA

(F) ZIP: 3000

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk

(B) COMPUTER: IBM PC compatible

(C) OPERATING SYSTEM: PC-DOS/MS-DOS

(D) SOFTWARE: PatentIn Release #1.0, Version #1.25

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: INTERNATIONAL (PCT)

(B) FILING DATE: 27-OCT-1999

(C) CLASSIFICATION:

(vi) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: AUSTRALIAN PROVISIONAL

(B) FILING DATE: 27-OCT-1998

(C) CLASSIFICATION:

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: HUGHES, DR E JOHN L

(C) REFERENCE/DOCKET NUMBER: EJH/AF

(ix) TELECOMMUNICATION INFORMATION:

- (A) TELEPHONE: +61 3 9254 2777
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(C) TELEX: AA 31787

(2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 27 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: Oligonucleotides

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

CGTGCTACTT CCATTTGTCA CGTCCTG

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(2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 21 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: Oligonucleotides

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

TGCCGTGATA GCAAATTTGA G

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(2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 22 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: Oligonucleotides

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

AGGAAGTGAG CATTATGGAT GA

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(2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 20 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: Oligonucleotides

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

TGCTCCTCGT GCCCAGCGTT

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(2) INFORMATION FOR SEQ ID NO:5:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 27 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: Oligonucleotides

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

TTAGAATTCC CCGAGGAGGA GCTGTAC

27

(2) INFORMATION FOR SEQ ID NO:6:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 27 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: Oligonucleotides

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

CTAGAATTCT GCAGTCCCAC CACCCCG

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